

1/7
POLYMORPHISMS IN THE TNFRSF11B GENE

| | | | | | |
|-----------------|------------|------------|----------------|-------------|------|
| ACAGCGAAC | CTAGAGCAA | GTGCCAA | ACT TCTGTCGATA | GCTTGAGGCT | |
| AGTGGAAAGA | CCTCGAGGAG | GCTACTCCAG | AAGTTCAGCG | CGTAGGAAGC | 100 |
| TCCGATACCA | ATAGCCCTT | GATGATGGTG | GGGTTGGTGA | AGGGAACAGT | |
| GCTCCGCAAG | GTTATCCCTG | CCCCAGGCAG | TCCAATTTC | ACTCTGCAGA | 200 |
| TTCTCTCTGG | CTCTAACTAC | CCCAGATAAC | AAGGAGTGAA | TGCAGAATAG | |
| CACGGGCTT | AGGGCCAATC | AGACATTAGT | TAGAAAAAATT | CCTACTACAT | 300 |
| GGTTTATGTA | AACTTGAAGA | TGAATGATTG | CGAACTCCCC | GAAAAGGGCT | |
| CAGACAATGC | CATGCATAAA | GAGGGGCCCT | GTAATTGAG | GTTCAGAAC | 400 |
| CCGAAGTGAA | GGGGTCAGGC | AGCCGGGTAC | GGCGGAAACT | CACAGCTTTC | |
| GCCCAGCGAG | AGGACAAAGG | TCTGGGACAC | ACTCCAAC | CGTCCGGATC | 500 |
| TTGGCTGGAT | CGGACTCTCA | GGGTGGAGGA | GACACAAGCA | CAGCAGCTGC | |
| T | | | | | |
| CCAGCGTGTG | CCCAGCCCTC | CCACCGCTGG | TCCCAGCTGC | CAGGAGGCTG | 600 |
| GCCGCTGGCG | GGAAAGGGCC | GGGAAACCTC | AGAGCCCCGC | GGAGACAGCA | |
| GCCGCCTTGT | TCCTCAGCCC | GGTGGCTTT | TTTCCCCTG | CTCTCCCAGG | 700 |
| GGACAGACAC | CACCGCCCCA | CCCCTCACGC | CCCACCTCCC | TGGGGGATCC | |
| T | | | | | |
| TTTCCGCCCC | AGCCCTGAAA | GCGTTAATCC | TGGAGCTTC | TGCACACCCC | 800 |
| C | | | | | |
| CCGACCGCTC | CCGCCAAGC | TTCCTAAAAA | AGAAAGGTGC | AAAGTTGGT | |
| CCAGGATAGA | AAAATGACTG | ATCAAAGGCA | GGCGATACTT | CCTGTTGCCG | 900 |
| GGACGCTATA | TATAACGTGA | TGAGCGCACG | GGCTGCGGAG | ACGCACCGGA | |
| GCGCTGCC | AGCCGCCGCC | TCCAAGCCCC | TGAGGTTTCC | GGGGACCAC | 1000 |
| ATGAACAAGT | TGCTGTGCTG | CGCGCTCGTG | GTAAGTCCCT | GGGCCAGCCG | |
| C | | | | | |
| T | | | | | |
| [exon 1: 1001.. | | | | | |
| ..1030] | | | | | |
| ACGGGTGCC | GGCGCCTGGG | GAGGCTGCTG | CCACCTGGTC | TCCCAACCTC | 1100 |
| CCAGCGGACC | GGCGGGGAGA | AGGCTCCACT | CGCTCCCTCC | CAGGAGAGGC | |
| A | | | | | |
| TTGGGGTTAG | GCTGGAGCAG | GAAACCGCTT | TCAAGTTATG | CCATGCTTCC | 1200 |
| CCTAGGGTGT | CCTTTACGC | TGCAAAGTTC | CTGCTGACTT | TATGGAAGAC | |
| A | | | | | |
| AGCAAGAGAG | AGACAGACAG | CGAGAGAGAG | GGAGAGAGAG | AGAGAGAGAA | 1300 |
| ACTTGTGTTGA | AAGTTTAGT | CATTAACCTT | CTGTCTTCAT | CTCAGAAATAT | |
| TAACGCCCTC | ATGTAGTCCA | TACTATCTT | GCTTAATGAA | CTTGAAC | 1400 |
| TATTATTAGT | GGCAAAGAAG | TGGTCCCTTA | GATTCAAGAGT | AAGTTGGAAG | |
| AAGACGTTAG | TCTTCTTAAA | ACCATTATAA | TTAGAATATG | ACATGATAGA | 1500 |
| NNNNNNNNNN | NNNNNNNNNN | NNNNNNNNNN | NNNNNNNNNN | NNNNNNNNNN | |
| CAGGACTTTG | AGTCAAATGA | TACTGTTGCA | CATAAGAAC | AACCTATTT | 1600 |
| CATGCTAAGA | TGATGCCACT | GTGTTCTT | CTCCTTCTAG | TTTCTGGACA | |
| [exon 2: 1641.. | | | | | |
| TCTCCATTAA | GTGGACCACC | CAGGAAACGT | TTCCCTCCAAA | GTACCTTCAT | 1700 |
| TATGACGAAG | AAACCTCTCA | TCAGCTGTTG | TGTGACAAAT | GTCCTCCTGG | |
| TACCTACCTA | AAACAACACT | GTACAGCAA | GTGGAAGACC | GTGTGCGCCC | 1800 |
| CTTGCCTGTA | CCACTACTAC | ACAGACAGCT | GGCACACCAG | TGACGAGTGT | |
| CTATACTGCA | GCCCCGTGTG | CAAGGAGCTG | CAGTACGTCA | AGCAGGAGTG | 1900 |
| CAATCGCACC | CACAACCGCG | TGTGCGAATG | CAAGGAAGGG | CGCTACCTTG | |
| AGATAGAGTT | CTGCTTGAAA | CATAGGAGCT | GCCCTCCTGG | ATTGGAGTG | 2000 |
| GTGCAAGCTG | GTACGTGTCA | ATGTGCAGCA | AAATTAATTA | GGATCATGCA | |
| T | | | | | |
| ..2010] | | | | | |

FIGURE 1A

| | | | | | |
|-------------|-------------|-------------|------------|-------------|------|
| AAGTCAGATA | GTTGTGACAG | TTTAGGAGAA | CACTTTGTT | CTGATGACAT | 2100 |
| TATAGGATAG | CAAATTGCAA | AGGTaatGAA | ACCTGCCAGG | TAGGTACTAT | |
| GTGTCTGGAG | TGCTTCCAAA | GGACCATTGC | TCAGAGGAAT | ACTTTGCCAC | 2200 |
| C | | | | | |
| TACAGGGCAA | TTTAATGACA | AATCTCAAAT | GCAGCAAATT | ATTCTCTCAT | |
| GAGATGCATG | ATGGTTTTT | TTTTTTTTT | TAAAGAAACA | AACTCAAGTT | 2300 |
| GCACATTGA | TAGTTGATCT | ATACCTCTAT | ATTCACCTTC | AGCAGGGACA | |
| CCTTCAAAC | GCAGCACTT | TTGACAAACA | TCAGAAATGT | TAATTTATAC | 2400 |
| CAAGAGAGTA | ATTATGCTCA | TATTAATGAG | ACTCTGGAGT | GCTAACAAATA | |
| AGCAGTTATA | ATTAATTATG | AAAAAAATGA | GAATGGTGAG | GGGAATTGCA | 2500 |
| TTTCATTATT | AAAAACAAAGG | CTAGTTCTTC | CTTAGCATG | GGAGCTGAGT | |
| GTGGGGAGG | GTAAGGACTA | TAGCAGAAC | TCTTCAATGA | GCTTATTCTT | 2600 |
| TATCTTAGAC | AAAACAGATT | GTCAAGCCAA | GAGCAAGCAC | TTGCCTATAA | |
| ACCAAGTGCT | TTCTCTTTG | CATTTGAAC | AGCATTGGTC | AGGGCTCATG | 2700 |
| TGTATTGAAT | CTTTAAACC | AGTAACCCAC | GTCCCCCTTC | TGCCACATTT | |
| GCGAAGCTTC | AGTGCAGCCT | ATAACTTTTC | ATAGCTTGAG | AAAATTAAGA | 2800 |
| GTATCCACTT | ACTTAGATGG | AAGAAGTAAT | CAGTATAGAT | TCTGATGACT | |
| CAGTTGAAG | CAGTGTTC | CAACTGAAGC | CCTGCTGATA | TTTTAAGAAA | 2900 |
| TATCTGGATT | CCTAGGCTGG | ACTCCTTTT | GTGGGCAGCT | GTCCTGCAGCA | |
| TTGTAGAATT | TTGGCAGCAC | CCCTGGACTC | TAGCCACTAG | ATACCAATAG | 3000 |
| CAGTCCTTCC | CCCATGTGAC | AGCCAAAAT | GTCTTCAGAC | ACTGTCAAAT | |
| GTCGCCAGGT | GGCAAAATCA | CTCCTGGTG | AGAACAGGGT | CATCAATGCT | 3100 |
| AAGTATCTGT | AACTATTTA | ACTCTAAAAA | CTTGTGATAT | ACAAAGTCTA | |
| AATTATTAGA | CGACCAATAC | TTAGGTTA | AAGGCATACA | AATGAAACAT | 3200 |
| TCAAAATCA | AAATCTATT | TGTTTCTCAA | ATAGTGAATC | TTATAAAATT | |
| AATCACAGAA | GATGCAAATT | GCATCAGAGT | CCCTAAAAT | TCCTCTCGT | 3300 |
| ATGAGTATT | GAGGGAGGAA | TTGGTGATAG | TTCCTACTT | CTATTGGATG | |
| GTACTTGAG | ACTCAAAAGC | TAAGCTAAGT | TGTGTGTGTG | TCAGGGTGC | 3400 |
| GGGTGTGGAA | TCCCATCAGA | AAAAAGCAA | TCCATGTAAT | TCATTCACTA | |
| AGTTGTATAT | GTAGAAAAAT | GAAAAGTGGG | CTATGCAGCT | TGGAAACTAG | 3500 |
| AGAATTTGA | AAAATAATGG | AAATCACAAG | GATCTTCTT | AAATAAGTAA | |
| GAAAATCTGT | TTGTAGAATG | AAGCAAGCAG | GCAGCCAGAA | GACTCAGAAC | 3600 |
| AAAAGTACAC | ATTTTACTCT | GTGTACACTG | GCAGCACAGT | GGGATTATT | |
| TACCTCTCCC | TCCCTAAAAA | CCCACACAGC | GGTCCTCTT | GGGAAATAAG | 3700 |
| AGGTTTCCAG | CCCAAAGAGA | AGGAAAGACT | ATGTGGTGTT | ACTCTAAAAA | |
| GTATTTAATA | ACCGTTTGT | TGTTGCTGTT | GCTGTTTGA | AATCAGATTG | 3800 |
| TCTCCTCTCC | ATATTTATT | TACTTCATT | TGTTAATTCC | TGTGGAATTA | |
| CTTAGAGCAA | GCATGGTGAA | TTCTCAACTG | TAAAGCCAAA | TTTCTCCATC | 3900 |
| ATTATAATT | CACATTTGC | CTGGCAGGTT | ATAATTTA | TATTTCCACT | |
| GATAGTAATA | AGGTAAAATC | ATTACTTAGA | TGGATAGATC | TTTTCATAA | 4000 |
| AAAGTACCAT | CAGTTATAGA | GGGAAGTCAT | GTTCATGTT | AGGAAGGTCA | |
| TTAGATAAAAG | CTTCTGAATA | TATTATGAAA | CATTAGTTCT | GTCATTCTTA | 4100 |
| GATTCTTTT | GTAAATAAC | TTTAAAAGCT | AACTTACCTA | AAAGAAATAT | |
| CTGACACATA | TGAACCTCTC | ATTAGGATGC | AGGAGAAGAC | CCAAGCCACA | 4200 |
| GATATGTATC | TGAAGAATGA | ACAAGATTCT | TAGGCCCGGC | ACGGTGGCTC | |
| ACATCTGTAA | TCTCAAGAGT | TTGAGAGGTC | AAGGCGGGCA | GATCACCTGA | 4300 |
| GGTCAGGAGT | TCAAGACCAG | CCTGGCCAAAC | ATGATGAAAC | CCTGCCTCTA | |
| CTAAAAATAC | AAAAATTAGC | AGGGCATGGT | GGTGCATGCC | TGCAACCTA | 4400 |
| GCTACTCAGG | AGGCTGAGAC | AGGAGAATCT | CTTGAACCT | CGAGGCAGGAG | |
| GTTGTGGTGA | GCTGAGATCC | CTCTACTGCA | CTCCAGCCTG | GGTGACAGAG | 4500 |
| ATGAGACTCC | GTCCCTGCCG | CCGCCCCCGC | CTTCCCCCCC | AAAAAGATTC | |
| TTCTTCATGC | AGAACATACG | GCAGTCAACA | AAGGGAGACC | TGGGTCCAGG | 4600 |
| TGTCCAAGTC | ACTTATTTCG | AGTAAATTAG | CAATGAAAGA | ATGCCATGGA | |
| ATCCCTGCC | AAATACCTCT | GCTTATGATA | TTGTAGAATT | TGATATAGAG | 4700 |

| | | |
|-----------------|--|------|
| TTGTATCCC | TTTAAGGAGT AGGATGAGT AGGAAAGTAC TAAAAACAAA | |
| CACACAAACA | GAAAACCCTC TTTGCTTG AAGGTGGTTC CTAAGATAAT | 4800 |
| GTCAGTGCAA | TGCTGGAAAT AATATTTAAT ATGTGAAGGT TTTAGGCTGT | |
| GTTTCCCT | CCTGTTCTT TTTCTGCCA GCCCTTGTC ATTGTTGCAG | 4900 |
| GTCAATGAAT | CATGTAGAAA GAGACAGGAG ATGAAACTAG AACCACTCCA | |
| TTTGCCCCCT | TTTTTATT TCTGGTTTG GTAAAAGATA CAATGAGGTA | 5000 |
| GGAGGTTGAG | ATTTATAAT GAAGTTAAT AAGTTCTGT AGCTTGATT | |
| TTTCTCTTC | ATATTGTTA TCTGCATAA GCCAGAATTG GCCTGTAAA | 5100 |
| TCTACATATG | GATATTGAAG TCTAAATCTG TTCAACTAGC TTACACTAGA | |
| TGGAGATATT | TTCATATTCA GATACACTGG AATGTATGAT CTAGCCATGC | 5200 |
| GTAATATAGT | CAAGTGTGTT AAGGTATT TA TTTTAATAG CGTCTTAGT | |
| TGTGGACTGG | TTCAAGTTT TCTGCCAATG ATTTCTCAA ATTATCAA | 5300 |
| TATTTTCCA | TCATGAAGTA AAATGCCCT GCAGTCACCC TTCTGAAGT | |
| TTGAACGACT | CTGCTGTTT AAACAGTTA AGCAAATGGT ATATCATCTT | 5400 |
| CCGTTACTA | TGTAGCTAA CTGCAGGCTT ACGCTTTGA GTCAGCGGCC | |
| AACTTTATTG | CCACCTCAA AAGTTTATT TAATGTTGTA AATTTTACT | 5500 |
| TCTCAAGGTT | AGCATACTTA GGAGTTGCTT CACAATTAGG ATTCAAGGAAA | |
| GAAAGAACTT | CAGTAGGAAC TGATTGGAAT TTAATGATGC AGCATTCAAT | 5600 |
| GGGTACTAAT | TTCAAAGAAT GATATTACAG CAGACACACA GCAGTTATCT | |
| TGATTTCTA | GGAATAATTG TATGAAGAAT ATGGCTGACA ACACGGCCTT | 5700 |
| ACTGCCACTC | AGCGGAGGCT GGACTAATGA ACACCCTACC CTTCTTCCT | |
| TTCCTCTCAC | ATTTCATGAG CGTTTGTAG GTAACGAGAA AATTGACTTG | 5800 |
| CATTTGCATT | ACAAGGAGGA GAAACTGGCA AAGGGGATGA TGGTGGAAAGT | |
| TTTGTCTGT | CTAATGAAGT GAAAAATGAA AATGCTAGAG TTTGTGCAA | 5900 |
| CATAATAGTA | GCAGTAAAAA CCAAGTGAAA AGTCTTCCA AACTGTGTT | |
| C | | |
| AAGAGGGCAT | CTGCTGGAA ACGATTGAG GAGAAGGTAC TAAATTGCTT | 6000 |
| GGTATTTCC | GTAGGAACCC CAGAGCGAAA TACAGTTGC AAAAGATGTC | |
| T | | |
| [exon 3: 6015.. | | |
| CAGATGGGTT | CTTCTCAAAT GAGACGTCA CAAAGCACC CTGTAGAAAA | 6100 |
| CACACAAATT | GCAGTGTCTT TGGTCTCCTG CTAACTCAGA AAGGAAATGC | |
| AACACACGAC | AACATATGTT CCGGAAACAG TGAATCAACT CAAAATGTG | 6200 |
| GAATAGGTAA | TTACATTCCA AAATACGTCT TTGTACGATT TTGTAGTATC | |
| ..6206] | | |
| ATCTCTCTCT | CTGAGTTGAA CACAAGGCCT CCAGCCACAT TCTTGGTCAA | 6300 |
| ACTTACATTT | TCCCTTTCTT GAATCTTAAC CAGCTAAGGC TACTCTCGAT | |
| GCATTACTGC | TAAAGCTACC ACTCAGAAC TCTCAAAAC TCATCTTCTC | 6400 |
| ACAGATAACA | CCTCAAAGCT TGATTTCTC TCCTTTCACA CTGAAATCAA | |
| ATCTTGCCCA | TAGGCAAAGG GCAGTGTCAA GTTGCCACT GAGATGAAAT | 6500 |
| TAGGAGAGTC | CAAACGTAG AATTACGTT GTGTGTTATT ACTTCACGA | |
| ATGTCTGTAT | TATTAACAA AGTATATATT GGCAACTAAG AAGCAAAGTG | 6600 |
| ATATAAACAT | GATGACAAAT TAGGCCAGGC ATGGTGGCTT ACTCCTATAA | |
| TCCCAACATT | TTGGGGGGCC AAGGTAGGCA GATCACTTGA GGTCAAGGATT | 6700 |
| TCAAGACCAAG | CCTGACCAAC ATGGTGAAC CTTGTCTCTA CTAAAATAC | |
| AAAAATTAGC | TGGGCATGGT AGCAGGCACT TCTAGTACCA GCTACTCAGG | 6800 |
| GCTGAGGCAG | GAGAATCGCT TGAACCCAGG AGATGGAGGT TGCAGTGAGC | |
| TGAGATTGTA | CCACTGCACT CCAGTCTGGG CAACAGAGCA AGATTCATC | 6900 |
| ACACACACAC | ACACACACAC ACACACACAC ATTAGAAATG TGTACTTGGC | |
| TTTGTACCT | ATGGTATTAG TGCATCTATT GCATGGAAC TCCAAGCTAC | 7000 |
| TCTGGTTGTG | TTAAGCTCTT CATGGGTAC AGGTCACTAG TATTAAGTTC | |
| AGGTTATTGCG | GATGCATTCC ACGGTAGTGA TGACAATTCA TCAGGCTAGT | 7100 |
| GTGTGTGTC | ACCTGTCAAC TCCCACCACT AGACTAATCT CAGACCTTCA | |
| CTCAAAGACA | CATTACACTA AAGATGATT GCTTTTGT GTTTAATCAA | 7200 |

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|--|--|
| GCAATGGTAT AAACCAGCTT GACTCTCCCC AAACAGTTT TCGTACTACA AAGAACGTTA TGAAGCAGAG AAATGTGAAT TGATATATAT ATGAGATTCT AACCCAGTTC CAGCATTGTT TCATTGTGTA ATTGAAATCA TAGACAAGCC ATTTTAGCCT TTGCTTCTT ATCTAAAAAA AAAAAAAA AAATGAAGGA AGGGGTATTA AAAGGAGTGA TCAAATTAA ACATTCTCTT TAATTAAATC ATTTTAATT TTACTTTTT TCATTATTG TGCACCTACT ATGTGGTACT GTGCTATAGA GGCTTAACA TTTATAAAA CACTGTGAAA GTTGCTTCAG ATGAATATAG GTAGTAGAAC GGCAGAACTA GTATTCAAAG CCAGGTCTGA TGAATCCAAA AACAAACACC CATACTCCC ATTTCTGGG ACATACTTAC TCTACCCAGA TGCTCTGGGC TTTGTAATGC CTATGAAAT AACATAGTT TATGTTGGT TATTTCCTA TGTAATGTCT ACTTATATAT CTGTATCTAT CTCTTGCTTT GTTCCAAAG GTAAACTATG TGTCTAAATG TGGGCAAAA ATAACACACT ATTCCAAATT ACTGTTCAAAT TTCCCTTAAG TCAGTGATAA TTATTTGTTT TGACATTAAT CATGAAGTTC CCTGTGGGTAA CTAGGTAAAC CTTTAATAGA ATGTTAATGT TTGTATTCTAT TATAAGAATT TTTGGCTGTT ACTTATTTAC AACAAATATT CACTCTAATT AGACATTTAC TAAACTTTCT CTTGAACACA ATGCCCAAAA AAGAACATTA GAAGACACGT AAGCTCAGTT GGTCTCTGCC ACTAAGACCA GCCAACAGAA GCTTGATTT ATTCAAACCTT TGCATTTAG CATATTTAT CTTGGAAAAT TCAATTGTGT TGGTTTTTG | 7300 7400 7500 7600 7700 7800 7900 8000 8100 |
| A | |
| TTTTGTTTG TATTGAATAG ACTCTCAGAA ATCCAATTGT TGAGTAAATC TTCTGGGTTT TCTAACCTTT CTTAGATGT TACCTGTGT GAGGAGGCAT [exon 4: 8227..] | 8200 |
| TCTTCAGGTT TGCTGTTCCCT ACAAAAGTTA CGCCTAACTG GCTTAGTGTC TTGGTAGACA ATTCGCCTGG CACCAAAGTA AACGCAGAGA GTGTAGAGAG | 8300 |
| T | |
| GATAAAACGG CAACACAGCT CACAAGAACAA GACTTTCCAG CTGCTGAAGT | 8400 |
| G | |
| TATGGAAACA TCAAAACAAA GACCAAGATA TAGTCAAGAA GATCATCCAA | |
| G | |
| GGTATGATAA TCTAAAATAA AAAGATCAAT CAGAAATCAA AGACACCTAT | 8500 |
| C | |
| ..8451] | |
| TTATCATAAA CCAGGAACAA GACTGCATGT ATGTTAGTT GTGTGGATCT TGTTCCCTG TTGGAATCAT TGTGGACTG AAAAAGTTTC CACCTGATAA TGTAGATGTG ATTCCACAAA CAGTTATACA AGGTTTGTT CTCACCCCTG CTCCCCAGTT TCCTGTAAA GTATGTTGAA CACTCTAAGA GAAGAGAAAT GCATTTGAAG GCAGGGCTGT ATCTCAGGGAA GTCGCTTCCA GATCCCTTAA CGCTCTGTAA AGCAGCCCCT CTAGACCACC AAGGAGAAC TCTATAACCA CTTTGTATCT TACATTGCAC CTCTACCAAG AAGCTCTGTT GTATTTACTT GGTAATTCTC TCCAGGTAGG CTTTCGTAG CTTACAAATA TGTTCTTATT AATCCTCATG ATATGGCCTG CATTAAAATT ATTTTAATGG CATATGTTAT GAGAATTAAT GAGATAAAAT CTGAAAAGTG TTTGAGCCTC TTGTAGGAAA AAGCTAGTTA CAGCAAAATG TTCTCACATC TTATAAGTTT ATATAAAGAT TCTCCTTCTAG AAATGGTGTG AGAGAGAAC AGAGAGAGAT AGGGAGAGAA GTGTGAAAGA ATCTGAAGAA AAGGAGTTTC ATCCAGTGTG GACTGTAAGC TTTACGACAC ATGATGGAAA GAGTTCTGAC TTCAGTAAGC ATTGGGAGGA CATGCTAGAA GAAAAAGGAA GAAGAGTTTC CATAATGCAG ACAGGGTCAG TGAGAAATTG ATTCAAGGTCC TCACCAAGTAG TTAAATGACT GTATAGTCTT GCACTACCCT AAAAAACTTC AAGTATCTGA AACCGGGGCA ACAGATTTA GGAGACCAAC GTCTTGAGA GCTGATTGCT TTTGCTTATG CAAAGAGTAA ACTTTTATGT TTTGAGCAAA CCAAAAGTAT TCTTTGAACG TATAATTAGC CCTGAAGCCG AAAGAAAAGA GAAAATCAGA GACCGTTAGA ATTGGAAGCA ACCAAATTCC CTATTTATAA AATGAGGACA TTTTAACCCA GAAAGATGAA | 8600 8700 8800 8900 9000 9100 9200 9300 9400 9500 |

| | | | | | |
|------------------|-------------|-------------|------------|-------------|-------|
| CCGATTGGC | TTAGGGCTCA | CAGATACTAA | GTGACTCATG | TCATTAATAG | 9600 |
| AAATGTTAGT | TCCTCCCTCT | TAGGTTGTA | CCCTAGCTA | TTACTGAAAT | |
| ATTCTCTAGG | CTGTGTGTCT | CCTTAGTTC | CTCGACCTCA | TGTCTTGAG | 9700 |
| TTTCAGATA | TCCTCCTCAT | GGAGGGTAGTC | CTCTGGTGCT | ATGTGTATT | |
| TTTAAAGGCT | AGTTACGGCA | ATTAACATTAT | CAACTAGCGC | CTACTAATGA | 9800 |
| AACTTTGTAT | TACAAAGTAG | CTAACTTGAA | TACTTTCTT | TTTTCTGAA | |
| ATGTTATGGT | GGTAATTCT | CAAACCTTTT | CTTAGAAAAC | TGAGAGTGAT | 9900 |
| GTGTCTTATT | TTCTACTGTT | AATTTCAAA | ATTAGGAGCT | TCTTCCAAAG | |
| TTTGTGGA | TGCCAAAAAT | ATATAGCATA | TTATCTTATT | ATAACAAAAAA | 10000 |
| ATATTATCT | CAGTTCTTAG | AAATAAATGG | TGTCACTTAA | CTCCCTCTCA | |
| AAAGAAAAGG | TTATCATTGA | AATATAATT | TGAAATTCTG | CAAGAACCTT | 10100 |
| TTGCCTCACG | CTTGTGTTAT | GATGGCATTG | GATGAATATA | AATGATGTGA | |
| ACACTTATCT | GGGCTTTGC | TTTATGCAGA | TATTGACCTC | TGTGAAAACA | 10200 |
| [exon 5: 10180.. | | | | | |
| GC GTGCAGCG | GCACATTGGA | CATGCTAAC | TCACCTTCGA | GCAGCTTCGT | |
| A | | | | | |
| AGCTTGATGG | AAAGCTTACC | GGGAAAGAAA | GTGGGAGCAG | AAGACATTGA | 10300 |
| AAAAACAATA | AAGGCATGCA | AACCCAGTGA | CCAGATCCTG | AAGCTGCTCA | |
| GT TTGTGGCG | AATAAAAAT | GGCGACCAAG | ACACCTTGAA | GGGCCTAATG | 10400 |
| CACGCACTAA | AGCACTCAA | GACGTACCA | TTTCCCAAA | CTGTCACTCA | |
| GAGTCTAAAG | AAGACCATCA | GGTCCTTCA | CAGCTTCACA | ATGTACAAAT | 10500 |
| TGTATCAGAA | GTTATTTTA | GAAATGATAG | GTAACCAGGT | CCAATCAGTA | |
| C | | | | | |
| AAAATAAGCT | GCTTATAAACT | GGAAATGGCC | ATTGAGCTGT | TTCCTCACAA | 10600 |
| ..10568] | | | | | |
| TTGGCGAGAT | CCCATGGATG | AGTAAACTGT | TTCTCAGGCA | CTTGAGGCTT | |
| TCAGTGATAT | CTTTCTCATT | ACCACTGACT | AATTTGCCA | CAGGGTACTA | 10700 |
| AAAGAAAACTA | TGATGTGGAG | AAAGGACTAA | CATCTCCTCC | AATAAACCCC | |
| AAATGGTTAA | TCCA ACTGTC | AGATCTGGAT | CGTTATCTAC | TGACTATATT | 10800 |
| TTCCCTTATT | ACTGCTTGCA | GTAATTCAAC | TGGAAATTAA | AAAAAAAAAAA | |
| CTAGACTCCA | CTGGGCCTTA | CTAAATATGG | GAATGTCTAA | CTTAAATAGC | 10900 |
| TTTGGGATTC | CAGCTATGCT | AGAGGCTTT | ATTAGAAAGC | CATATTTTT | |
| TCTGTAAAAG | TTACTAATAT | ATCTGTAACA | CTATTACAGT | ATTGCTATT | 11000 |
| ATATTCAATT | AGATATAAGA | TTTGGACATA | TTATCATCCT | ATAAAGAAC | |
| GGTATGACTT | AATTTAGAA | AGAAAATTAT | ATTCTGTTA | TTATGACAAA | 11100 |
| TGAAAGAGAA | AATATATATT | TTAATGGAA | AGTTTGTAGC | ATTTTCTAA | |
| TAGGTACTGC | CATATTTTC | TGTGTGGAGT | ATTTTATAA | TTTTATCTGT | 11200 |
| ATAAGCTGTA | ATATCATT | ATAGAAAATG | CATTATTAG | TCAATTGTT | |
| AATGTTGGAA | AACATATGAA | ATATAAATT | TCTGAATATT | AGATGCTCTG | 11300 |
| AGAAATTGAA | TGTACCTTAT | TTAAAAGATT | TTATGGTTT | ATAACTATAT | |
| AAATGACATT | ATTAAAGTTT | TCAAATTATT | TTTATTGCT | TTCTCTGTTG | 11400 |
| CTTTTATT | | | | | 11408 |

POLYMORPHISMS IN THE CODING SEQUENCE OF TNFRSF11B

| | | | | | |
|-------------|-------------|-------------|------------|-------------|------|
| ATGAACAAAGT | TGCTGTGCTG | CGCGCTCGTG | TTTCTGGACA | TCTCCATTAA | |
| C | | | | | |
| GTGGACCACC | CAGGAAACGT | TTCCTCCAAA | GTACCTTCAT | TATGACGAAG | 100 |
| AAACCTCTCA | TCAGCTGTTG | TGTGACAAAT | GTCCTCCTGG | TACCTACCTA | |
| AAACAAACACT | GTACAGCAAA | GTGGAAGACC | GTGTGCGCCC | CTTGCCTGAA | 200 |
| CCACTACTAC | ACAGACAGCT | GGCACACCAAG | TGACGAGTGT | CTATACTGCA | |
| GCCCCGTGTG | CAAGGAGCTG | CAGTACGTCA | AGCAGGAGTG | CAATCGCACC | 300 |
| CACAACCGCG | TGTGCGAATG | CAAGGAAGGG | CGCTACCTTG | AGATAGAGTT | |
| CTGCTTGAAA | CATAGGAGCT | GCCCTCCTGG | ATTGGAGTG | GTGCAAGCTG | 400 |
| GAACCCCAGA | GCGAAATACA | GTTTGCAAAA | GATGTCCAGA | TGGGTTCTTC | |
| TCAAATGAGA | CGTCATCTAA | AGCACCCGT | AGAAAACACA | CAAATTGCAG | 500 |
| TGTCTTGTT | CTCCTGCTAA | CTCAGAAAGG | AAATGCAACA | CACGACAACA | |
| TATGTTCCGG | AAACAGTGAA | TCAACTCAAA | AATGTGGAAT | AGATGTTACC | 600 |
| CTGTGTGAGG | AGGCATTCTT | CAGGTTGCT | GTTCCTACAA | AGTTTACGCC | |
| TAACTGGCTT | AGTGTCTTGG | TAGACAATT | GCCTGGCACC | AAAGTAAACG | 700 |
| T | | | | | |
| CAGAGAGTGT | AGAGAGGATA | AAACGGCAAC | ACAGCTCACA | AGAACAGACT | |
| G | | | | | |
| TTCCAGCTGC | TGAAGTTATG | GAAACATCAA | AACAAAGACC | AAGATATAGT | 800 |
| G | | | | | |
| CAAGAAGATC | ATCCAAGATA | TTGACCTCTG | TGAAAACAGC | GTGCAGCGGC | |
| A | | | | | |
| ACATTGGACA | TGCTAACCTC | ACCTTCGAGC | AGCTTCGTAG | CTTGATGGAA | 900 |
| AGCTTACCGG | GAAAGAAAGT | GGGAGCAGAA | GACATTGAAA | AAACAATAAA | |
| GGCATGCAA | CCCAGTGACC | AGATCCTGAA | GCTGCTCAGT | TTGTGGCGAA | 1000 |
| TAAAAAAATGG | CGACCAAGAC | ACCTTGAGG | GCCTAATGCA | CGCACTAAAG | |
| CACTCAAAGA | CGTACCACTT | TCCCAAAACT | GTCACTCAGA | GTCTAAAGAA | 1100 |
| GACCATCAGG | TTCCCTTCACA | GCTTCACAAT | GTACAAATTG | TATCAGAAAGT | |
| C | | | | | |
| TATTTTTAGA | AATGATAGGT | AACCAGGTCC | AATCAGTAAA | AATAAGCTGC | 1200 |
| TTATAA | | | | | 1206 |

ISOFORMS OF THE TNFRSF11B PROTEIN

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|---|-----|
| MNKLLCCALV FLDISIKWTT QETFPPKYLH YDEETSHQLL CDKCPPGTYL | |
| N | |
| KQHCTAKWKT VCAPCPDHYY TDSWHTSDEC LYCSPVCKEL QYVKQECNRT | 100 |
| HNRVCECKEG RYLEIEFCLK HRSCPFGV VQAGTPERNNT VCKRCPDGFF | |
| SNETSSKAPC RKHTNCVFG LLLTQGNAT HDNICSGNSE STQKCGIDVT | 200 |
| LCEEAFFRFA VPTKFTPWL SVLVDNLPGT KVNAESVERI KRQHSSQEQT | |
| M | |
| FQLLKLWKHQ NKDQDIVKKI IQDIDLCE NS VQRHIGHANL TFEQLRSLME | 300 |
| M | |
| SLPGKKVGAE DIEKTIKACK PSDQILKLLS LWRIKNGDQD TLKGLMHALK | 400 |
| HSKYHFPKT VTQSLKKTIR FLHSFTMYKL YQKLFLEMIG NQVQSVKISC | |
| L | 401 |